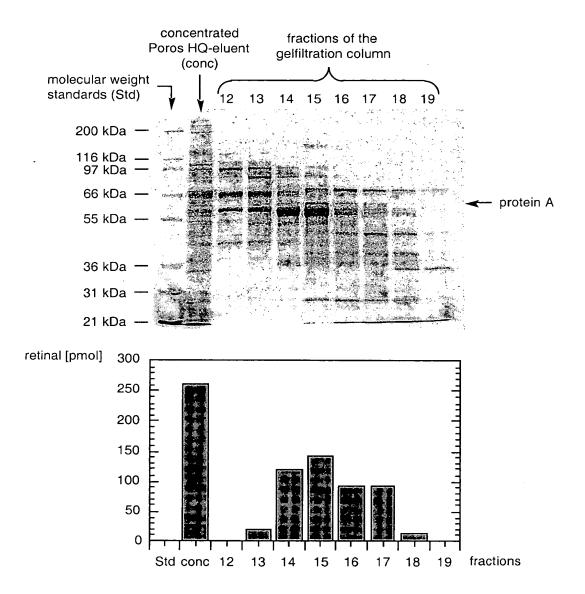
Figure 1



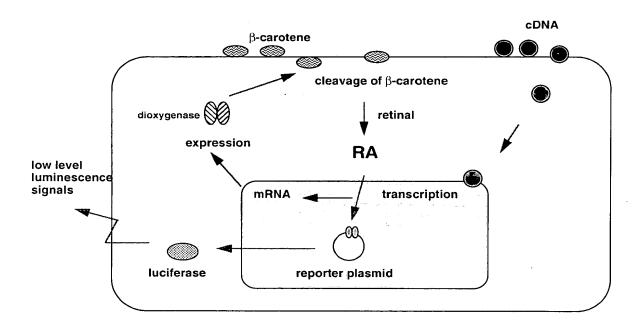


Figure 2

Figure 3 No. 2

Seq. ID

- 1 CGGATCCACT AGTAACGGCC GCCAGTGTGG TGGAATCCAT
- 51 AACAGGAAAG AGCTGTTCTT AGCCCAGAGA GGAGGGCACC GTACGCCTGC
- 101 AGGAGCAGCT GGGTAGAGGA CACAGGAGAG CGATGGAGAC AATATTTAAC
- 151 AGAAACAAAG AAGAGCATCC AGAGCCCATA AAAGCTGAGG TGCAAGGTCA
- 201 GTTGCCCACT TGGTTGCAAG GGGTACTTCT CCGAAATGGC CCAGGGATGC
- 251 ACACAATAGG GGACACTAAA TACAACCACT GGTTTGATGG
- 301 CTGCACAGCT TCACGTTTAA AAATGGTGAA GTTTACTACA GAAGTAAGTA
- 351 CCTCCGAAGT GACACATACA ACTGCAATAT AGAAGCAAAC CGAATCGTGG
- 401 TGTCTGAGTT TGGAACCATG GCTTATCCGG ATCCATGCAA AAACATATTT
- 451 GCCAAGGCAT TCTCATACTT ATCTCACACC ATTCCTGAGT TCACGGACAA
- 501 CTGCCTGATC AACATTATGA AAACTGGGGA TGATTATTAT GCTACCAGTG
- 551 AGACTAACTT CATCAGAAAA ATTGATCCAC AGACTCTGGA GACACTAGAT
- 601 AAGGTAGACT ACAGCAAATA TGTAGCTGTA AACTTGGCAA
- 651 ACACTATGAC AGTGCTGGAA ATATTCTCAA CATGGGTACT
- 701 ATAAAGGAG AACAAAATAT GTTCTCTTTA AGATCCCTTCCTTGTACCA
- 751 GAAAAAGAAA AGAAGAAATC TTGTTTTAAA CACCTGGAAG TAGTATGCTC
- 801 CATCCCTTCT CGCTCCCTGC TCCAACCAAG CTACTACCAC AGCTTTGGAA
- 851 TCACAGAAAA TTATATTGTG TTCATAGAGC AGCCATTTAA ACTGGATATT

- 901 GTCAAACTGG CAACTGCCTA CATCCGAGGT GTGAACTGGG
- 951 TTCCTTTCAT AAGGAGGATA AGACGTGGTT TCACTTTGTA GACAGAAGA
- 1001 CGAAAAAGA AGTATCCAÇC AAGTTTTACA CTGATGCTTT GGTGCTTTAT
- 1051 CACCACATAA ATGCTTACGA AGAAGATGGC CACGTTGTTT TTGATATCGT
- 1101 TGCCTACAGA GACAATAGCT TGTACGATAT GTTTTACTTA AAAAAACTGG
- 1151 ACAAAGACTT TGAAGTGAAC AACAAGCTTA CCTCCATCCC AACCTGCAAG
- 1201 CGCTTTGTTG TGCCTCTGCA GTATGACAAG GATGCAGAAG TAGGTTCTAA
- 1251 TTTAGTCAAA CTTCCAACTT CCGCAACTGC TGTAAAAGAA AAAGATGGCA
- 1301 GCATCTATTG TCAACCTGAA ATATTATGTG AAGGGATAGA ACTGCCTCGT
- 1351 GTCAACTATG ACTACAATGG CAAAAAATAC AAGTATGTCT ATGCAACAGA
- 1401 AGTCCAGTGG AGCCCAGTTC CTACAAAGAT TGCAAAACTG AATGTCCAAA
- 1451 CAAAGGAAGT ACTGCACTGG GGAGAAGACC ACTGCTGGCC CTCAGAGCCC
- 1501 ATCTTTGTTC CCAGCCCCGA TGCAAGAGAA GAGGATGAAG GTGTTGTTTT
- 1551 GACCTGTGTT GTGGTGTCTG AGCCAAATAA AGCACCCTTC CTACTCATCT
- 1601 TGGATGCTAA AACATTCAAA GAATTGGGCC GAGCCACAGT TAACGTAGAA
- 1651 ATGCATCTGG ACCTGCATGG GATGTTTATA CCACAGAATG ATTTGGGGGC
- 1701 TGAGACGGAA TAAAACGCTA TTGATCCGAC TACACAAACT GAGACAACTT
- 1751 TCTACTGAAC ATGAGTTAAT ATCCCTTTTA CCATTCAAGA ACAACCATAT
- 1801 AACGACACAA AATGACTATG TATAATCTCT TAAATAATAG ATATAATCCT
- 1851 TTTAAGGCAC AGCGATGAGT TTTACTACAG GTAACGATAT GCACAACTGG

- 1901 CATATAACTA TTCCAAAAGA AGAAGAACGA TCAGTGTTTT AGAAGTGCTA
- 1951 ATGTTGTACA TAACGGCGGC AGAGGGAACA GGAGAGAAG GTAACGGGAA
- 2001 TATTTAATAG AATATAGATT TCTGAGCAAA TGAAGTGCAG TATTTATGGT
- 2051 GTGATGCATG GCATGAGTCA CATAGGTCTG CAGCTCATGT ATCTTTAGA
- 2101 GATCGTTTCA AGATTGCAGC TTGTGATGCA AGTTTTCTCC AGCCAGAAAA
- 2151 CCTCATTTTA AACCATCTGC TACTGGTAAT TCATACCAAT GCATTTTCTT
- 2201 GGTGCTCGAT TTACACTATA ACCAAAGTTA AGTATTACAT TCAGGTGCTA
- 2251 CAACTTTCTA ATTTACAACC GAAACAAACA AGCAAACAGC ACTTGCTTTG
- 2301 CTAATAACCC CATGGTGTAT TTTTCCTTTT TATGATGACA AAACCAAGTA
- 2351 CATATGGTTT TATGTAGCAT TCAATTATAC TTCAGTGCTA
- 2401 ATGTTATAAG CAATTTGTAT TTAAATCAGT TTTCCTTGAG AATATCTGAC
- 2451 ATAACATTTT GTGTAATGAG ATGACTATGT TGTCTAAAGA TGAACAGGAA
- 2501 TGTATCTTTT ATTAGTATTG TTAATTGTGT TACTAATACT ATGCATATGA
- 2551 ATGAGAGCAA TGTATTTCTA GGAGAACTCA GATATACATT CAACAATTTC
- 2601 TGTAGGTGAA AATGCATTTA CTGATGAAAG TTGAATCGTT AATGAGGGAG
- 2651 AAAACTGGGT ATCCATCCAT CCAACTATGT TAGGTGTTCA CCTGGTCTGT
- 2701 ATGTGACACC ACGCTGTTTG GGTATCTCTC ACTTTCACAT ACCTGTTCTC
- 2751 ATGGTTTCTG CTACTCACTG TATTTTGCAG GAGAGAAACA AAATGAAATC
- 2801 ACTGTCACTT ACTATCGCCC CATCACATAA GAACAATGGG GCTTTGGTGA

- 2851 CTTGTTCATG ATTACATAAG ATGTTTGCAG CAGAGCAGCA ATAGAACCAA
- 2901 CACCATCCAC AGTTCTTGCT TGCTCTGTTA TGACTCCCTT TGCTGTCTTT
- 2951 ATGGTTTGCA TGTATGAAGA ATACACTGCC TAATTCTAAT GTTAAAAAGT
- 3001 CACTGGGGTC AGATCTAGAG CTTAAGTAAG CAGTCTGGGG TTTTCAAATG
- 3051 TTTATATGTT CCATAAAATG GAAATAAACA CCTCCATAAT AAAAAAAAA
 - 3101 AAAAAAAAA A

Figure 4 No. 1

Seq. ID

- 1 METIFNRNKE EHPEPIKAEV QGQLPTWLQG VLLRNGPGMH
- 51 FDGLALLHSF TFKNGEVYYR SKYLRSDTYN CNIEANRIVV SEFGTMAYPD
- 101 PCKNIFAKAF SYLSHTIPEF TDNCLINIMK TGDDYYATSE TNFIRKIDPO
- 151 TLETLDKVDY SKYVAVNLAT SHPHYDSAGN ILNMGTSIVD KGRTKYVLFK
- 201 IPSSVPEKEK KKSCFKHLEV VCSIPSRSLL QPSYYHSFGI TENYIVFIEQ
- 251 PFKLDIVKLA TAYIRGVNWA SCLSFHKEDK TWFHFVDRKT KKEVSTKFYT
- 301 DALVLYHHIN AYEEDGHVVF DIVAYRDNSL YDMFYLKKLD KDFEVNNKLT
- 351 SIPTCKRFVV PLQYDKDAEV GSNLVKLPTS ATAVKEKDGS IYCQPEILCE
- 401 GIELPRVNYD YNGKKYKYVY ATEVQWSPVP TKIAKLNVQT KEVLHWGEDH
- 451 CWPSEPIFVP SPDAREEDEG VVLTCVVVSE PNKAPFLLIL DAKTFKELGR
 - 501 ATVNVEMHLD LHGMFIPQND LGAETE

Figure 5		Seq ID No. 4 and Seq ID No. 5
57	10	EEHPEPIKAEVQGQLPTWLQGVLLRNGPGMHTIGDTKYNHWFDGLALL
69	20	: .: : : : EELSSPLTAHVTGRIPLWLTGSLLRCFTGPGLFEVGSEPFYHLFDGQALL
105	58	
119	70	
155	106	
167	120	.: : : : . . : : FSRFFSYFRGVEVTDNALVNVYPVGEDYYACTETNFITKINPETLETI
203	156	DKVDYSKYVAVNLATSHPHYDSAGNILNMGTSIVDKGRTKYVLFKIPS
217	168	. . . :. : . : FTKQVDLCNYVSVNGATAHPHIENDGTVYNIGNCFGKNFSIAYNIVKIPP
253	204	SVPEKEKKKSCFKHLEVVCSIPSRSLLQPSYYHSFGITENYIVFIEQPFK
	218	: : . : : LQADKEDPISKFTS.EIVVQFPCSDRFKPSYVHSFGLTPNYIVFVETPVK
266	254	
300	267	:.: : : : : : INLFKFLSSWSLWGANYMDCFESFTNETMGVWLHIADKKRKKYLNNKYRT
316	301	
344	317	: :. .: : :: SPFNLFHHINTYEDNGFLIVDLCCWKGFEFVYNYFTLYLANLRENWEEVK
366	345	. VNNKLTSIPTCKRFVVPLQYDKDAEVGSNLVKLP.TSATAVKEKDGSI
391	367	:
415	392	YCQPEILCEGIELPRVNYD.YNGKKYKYVYATEVQWSPVPTKIAKL
436		: : : :: WLEPEVLFSGPRQAFEFPQINYQKYCGKPYTYAYGLGLNHF.VPDRLCKL
464	410	WEBLEVEL SOLINGEL ELL ÉTIVI ÉTIVI CONFILITAT GEGENTIT VEDNECKE

484	43/	NVQTKEVLHWGEDHCWPSEPIFVPSPDAREEDEGVVLTCVVVSEPNKA
		. : : . NVKTKETWFTVWQEPDSYPSEPIFVSHPDALEEDDGVVLSVVVSPGAGQK
514	465	NVKTKETWFTVWQEPDSYPSEPIFVSHPDALEEDDGVVLSVVVSPGAGQK
	485	P.FLLILDAKTFKELGRATVNVEMHLDLHGMF 515
	515	: . : :

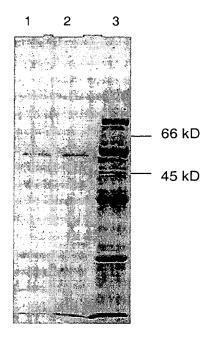


Fig. 6 shows a 10% polyacrylamide gel with E.coli expressed β,β -carotene 15,15'-dioxygenase after affinity tag purification; lane 1 and lane 2: 2 fractions from the Co^2 -chelate column showing the main band at 60 kD; lane 3: low range molecular weight marker (Bio Rad).

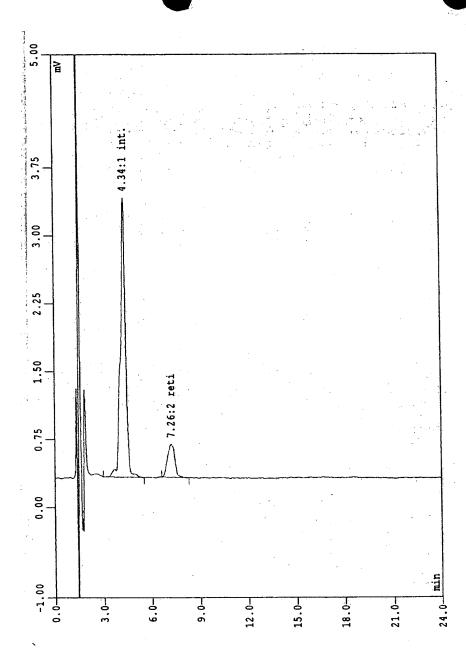


Fig. 7 shows an HPLC profile of the reaction mixture at the end of an activity assay for the β,β -carotene 15,15'-dioxygenase following the procedure in example 1. The first peak in the chromatogram represents the internal standard, while the second peak corresponds to retinal as the only product formed during the central cleavage with β -carotene as substrate.

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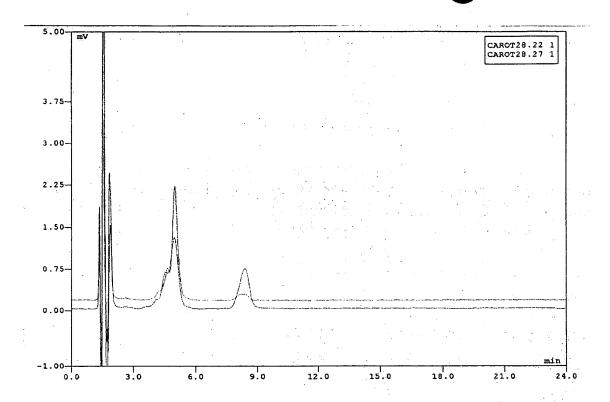


Fig. 8 confirms that the product peak in Fig. 7 is indeed retinal. A sample which was positive in the activity assay (green chromatogram) was spiked with retinal and analysed in second HPLC run (red chromatogram). The chromatograms of the two runs were then overlayed.